RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: _10/537,512_
Source: _PY7/10_
Date Processed by STIC: _6/14/05_

ENTERED

PCT

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    3 <110> APPLICANT: LYNCH, Berkley
    4        NOCKA, Karl
    5        FUKS, Bruno
    7 <120> TITLE OF INVENTION: Methods for the identification of agents for the treatment
of
    8        seizures, neurological diseases,endocrinopathies and hormonal
    9        diseases
   11 <130> FILE REFERENCE: 53529-5007-02-US
C--> 13 <140> CURRENT APPLICATION NUMBER: US/10/537,512
C--> 13 <141> CURRENT FILING DATE: 2005-06-03
   13 <150> PRIOR APPLICATION NUMBER: PCT/US2003/038122
   14 <151> PRIOR FILING DATE: 2003-12-02
   16 <150> PRIOR APPLICATION NUMBER: US 60/430,372
   17 <151> PRIOR FILING DATE: 2002-12-03
   19 <150> PRIOR APPLICATION NUMBER: US 60/506,764
   20 <151> PRIOR FILING DATE: 2003-09-30
   22 <160> NUMBER OF SEQ ID NOS: 16
   24 <170> SOFTWARE: PatentIn version 3.3
   26 <210> SEQ ID NO: 1
   27 <211> LENGTH: 2229
   28 <212> TYPE: DNA
   29 <213> ORGANISM: Homo sapiens
   32 <220> FEATURE:
   33 <221> NAME/KEY: CDS
   34 <222> LOCATION: (1)..(2229)
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   39 1               5                  10                  15
   41 gac att gct aag gaa gtc aaa aag cat gcg gcc aag aag gtg gtg aag     96
   42 Asp Ile Ala Lys Glu Val Lys Lys His Ala Ala Lys Lys Val Val Lys
   43             20                  25                  30
   45 ggc ctg gac aga gtc cag gac gaa tat tcc cga aga tcg tac tcc cgc    144
   46 Gly Leu Asp Arg Val Gln Asp Glu Tyr Ser Arg Arg Ser Tyr Ser Arg
   47         35                  40                  45
   49 ttt gag gag gag gat gat gat gat gac ttc cct gct ccc agt gat ggt    192
   50 Phe Glu Glu Glu Asp Asp Asp Asp Asp Phe Pro Ala Pro Ser Asp Gly
   51     50                  55                  60
   53 tat tac cga gga gaa ggg acc cag gat gag gag gaa ggt ggt gca tcc    240
   54 Tyr Tyr Arg Gly Glu Gly Thr Gln Asp Glu Glu Glu Gly Gly Ala Ser
   55 65                  70                  75                  80
   57 agt gat gct act gag ggc cat gac gag gat gat gag atc tat gaa ggg    288
   58 Ser Asp Ala Thr Glu Gly His Asp Glu Asp Asp Glu Ile Tyr Glu Gly
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RAW SEQUENCE LISTING   DATE: 06/14/2005
PATENT APPLICATION: **US/10/537,512**   TIME: 09:05:02

Input Set : **A:\53529-5007-02-US Sequence Listing.txt**
Output Set: **N:\CRF4\06142005\J537512.raw**

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 62 Glu Tyr Gln Gly Ile Pro Arg Ala Glu Ser Gly Gly Lys Gly Glu Arg
 63             100                 105                 110
 65 atg gca gat ggg gcg ccc ctg gct gga gta agg ggg ggc ttg agt gat        384
 66 Met Ala Asp Gly Ala Pro Leu Ala Gly Val Arg Gly Gly Leu Ser Asp
 67             115                 120                 125
 69 ggg gag ggt ccc cct ggg ggc cgg ggg gag gca caa cga cgg aaa gaa        432
 70 Gly Glu Gly Pro Pro Gly Gly Arg Gly Glu Ala Gln Arg Arg Lys Glu
 71      130                 135                 140
 73 cga gaa gaa ctg gcc caa cag tat gaa gcc atc cta cgg gag tgt ggc        480
 74 Arg Glu Glu Leu Ala Gln Gln Tyr Glu Ala Ile Leu Arg Glu Cys Gly
 75 145                 150                 155                 160
 77 cac ggc cgc ttc cag tgg aca ctg tat ttt gtg ctt ggt ctg gcg ctg        528
 78 His Gly Arg Phe Gln Trp Thr Leu Tyr Phe Val Leu Gly Leu Ala Leu
 79                 165                 170                 175
 81 atg gct gac ggt gtg gag gtc ttt gtg gtg ggc ttc gtg ctg ccc agc        576
 82 Met Ala Asp Gly Val Glu Val Phe Val Val Gly Phe Val Leu Pro Ser
 83             180                 185                 190
 85 gct gag aaa gac atg tgc ctg tcc gac tcc aac aaa ggc atg cta ggc        624
 86 Ala Glu Lys Asp Met Cys Leu Ser Asp Ser Asn Lys Gly Met Leu Gly
 87             195                 200                 205
 89 ctc atc gtc tac ctg ggc atg atg gtg gga gcc ttc ctc tgg gga ggt        672
 90 Leu Ile Val Tyr Leu Gly Met Met Val Gly Ala Phe Leu Trp Gly Gly
 91      210                 215                 220
 93 ctg gct gac cgg ctg ggt cgg agg cag tgt ctg ctc atc tcg ctc tca        720
 94 Leu Ala Asp Arg Leu Gly Arg Arg Gln Cys Leu Leu Ile Ser Leu Ser
 95 225                 230                 235                 240
 97 gtc aac agc gtc ttc gcc ttc ttc tca tct ttt gtc cag ggt tac ggc        768
 98 Val Asn Ser Val Phe Ala Phe Phe Ser Ser Phe Val Gln Gly Tyr Gly
 99                 245                 250                 255
101 act ttc ctc ttc tgc cgc cta ctt tct ggg gtt ggg att gga ggg tcc        816
102 Thr Phe Leu Phe Cys Arg Leu Leu Ser Gly Val Gly Ile Gly Gly Ser
103                 260                 265                 270
105 atc ccc att gtc ttc tcc tat ttc tcc gag ttt ctg gcc cag gag aaa        864
106 Ile Pro Ile Val Phe Ser Tyr Phe Ser Glu Phe Leu Ala Gln Glu Lys
107             275                 280                 285
109 cga ggg gag cat ttg agc tgg ctc tgc atg ttt tgg atg att ggt ggc        912
110 Arg Gly Glu His Leu Ser Trp Leu Cys Met Phe Trp Met Ile Gly Gly
111      290                 295                 300
113 gtg tac gca gct gct atg gcc tgg gcc atc atc ccc cac tat ggg tgg        960
114 Val Tyr Ala Ala Ala Met Ala Trp Ala Ile Ile Pro His Tyr Gly Trp
115 305                 310                 315                 320
117 agt ttt cag atg ggt tct gcc tac cag ttc cac agc tgg agg gtc ttc       1008
118 Ser Phe Gln Met Gly Ser Ala Tyr Gln Phe His Ser Trp Arg Val Phe
119                 325                 330                 335
121 gtc ctc gtc tgc gcc ttt cct tct gtg ttt gcc att ggg gct ctg acc       1056
122 Val Leu Val Cys Ala Phe Pro Ser Val Phe Ala Ile Gly Ala Leu Thr
123                 340                 345                 350
125 acg cag cct gag agc ccc cgt ttc ttc cta gag aat gga aag cat gat       1104
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126 Thr Gln Pro Glu Ser Pro Arg Phe Phe Leu Glu Asn Gly Lys His Asp
127         355             360             365
129 gag gcc tgg atg gtg ctg aag cag gtc cat gat acc aac atg cga gcc   1152
130 Glu Ala Trp Met Val Leu Lys Gln Val His Asp Thr Asn Met Arg Ala
131         370             375             380
133 aaa gga cat cct gag cga gtg ttc tca gta acc cac att aag acg att   1200
134 Lys Gly His Pro Glu Arg Val Phe Ser Val Thr His Ile Lys Thr Ile
135 385             390             395             400
137 cat cag gag gat gaa ttg att gag atc cag tcg gac aca ggg acc tgg   1248
138 His Gln Glu Asp Glu Leu Ile Glu Ile Gln Ser Asp Thr Gly Thr Trp
139             405             410             415
141 tac cag cgc tgg ggg gtc cgg gcc ttg agc cta ggg ggg cag gtt tgg   1296
142 Tyr Gln Arg Trp Gly Val Arg Ala Leu Ser Leu Gly Gly Gln Val Trp
143         420             425             430
145 ggg aat ttt ctc tcc tgt ttt ggt ccc gaa tat cgg cgc atc act ctg   1344
146 Gly Asn Phe Leu Ser Cys Phe Gly Pro Glu Tyr Arg Arg Ile Thr Leu
147         435             440             445
149 atg atg atg ggt gtg tgg ttc acc atg tca ttc agc tac tat ggc ctg   1392
150 Met Met Met Gly Val Trp Phe Thr Met Ser Phe Ser Tyr Tyr Gly Leu
151         450             455             460
153 acc gtc tgg ttt cct gac atg atc cgc cat ctc cag gca gtg gac tac   1440
154 Thr Val Trp Phe Pro Asp Met Ile Arg His Leu Gln Ala Val Asp Tyr
155 465             470             475             480
157 gca tcc cgc acc aaa gtg ttc ccc ggg gag cgc gta gag cat gta act   1488
158 Ala Ser Arg Thr Lys Val Phe Pro Gly Glu Arg Val Glu His Val Thr
159         485             490             495
161 ttt aac ttc acg ttg gag aat cag atc cac cga ggc ggg cag tac ttc   1536
162 Phe Asn Phe Thr Leu Glu Asn Gln Ile His Arg Gly Gly Gln Tyr Phe
163         500             505             510
165 aat gac aag ttc att ggg ctg cgg ctc aag tca gtg tcc ttt gag gat   1584
166 Asn Asp Lys Phe Ile Gly Leu Arg Leu Lys Ser Val Ser Phe Glu Asp
167         515             520             525
169 tcc ctg ttt gaa gag tgt tat ttt gag gat gtc aca tcc agc aac acg   1632
170 Ser Leu Phe Glu Glu Cys Tyr Phe Glu Asp Val Thr Ser Ser Asn Thr
171         530             535             540
173 ttt ttc cgc aac tgc aca ttc atc aac act gtg ttc tat aac act gac   1680
174 Phe Phe Arg Asn Cys Thr Phe Ile Asn Thr Val Phe Tyr Asn Thr Asp
175 545             550             555             560
177 ctg ttc gag tac aag ttt gtg aac agc cgt ctg ata aac agt aca ttc   1728
178 Leu Phe Glu Tyr Lys Phe Val Asn Ser Arg Leu Ile Asn Ser Thr Phe
179             565             570             575
181 ctg cac aac aag gag ggc tgc ccg cta gac gtg aca ggg acg ggc gaa   1776
182 Leu His Asn Lys Glu Gly Cys Pro Leu Asp Val Thr Gly Thr Gly Glu
183         580             585             590
185 ggt gcc tac atg gta tac ttt gtg agc ttc ctg ggg aca ctg gca gtg   1824
186 Gly Ala Tyr Met Val Tyr Phe Val Ser Phe Leu Gly Thr Leu Ala Val
187         595             600             605
189 ctt cct ggg aat atc gtg tct gcc ctg ctc atg gac aag atc ggc agg   1872
190 Leu Pro Gly Asn Ile Val Ser Ala Leu Leu Met Asp Lys Ile Gly Arg
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                    RAW SEQUENCE LISTING              DATE: 06/14/2005
                    PATENT APPLICATION:  US/10/537,512  TIME: 09:05:02

                    Input Set : A:\53529-5007-02-US Sequence Listing.txt
                    Output Set: N:\CRF4\06142005\J537512.raw
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194 Leu Arg Met Leu Ala Gly Ser Ser Val Met Ser Cys Val Ser Cys Phe
195 625                    630                    635                    640
197 ttc ctg tct ttt ggg aac agt gag tcg gcc atg atc gct ctg ctc tgc      1968
198 Phe Leu Ser Phe Gly Asn Ser Glu Ser Ala Met Ile Ala Leu Leu Cys
199                    645                    650                    655
201 ctt ttt ggc ggg gtc agc att gca tcc tgg aat gcg ctg gac gtg ttg      2016
202 Leu Phe Gly Gly Val Ser Ile Ala Ser Trp Asn Ala Leu Asp Val Leu
203                    660                    665                    670
205 act gtt gaa ctc tac ccc tca gac aag agg acc aca gct ttt ggc ttc      2064
206 Thr Val Glu Leu Tyr Pro Ser Asp Lys Arg Thr Thr Ala Phe Gly Phe
207            675                    680                    685
209 ctg aat gcc ctg tgt aag ctg gca gct gtg ctg ggg atc agc atc ttc      2112
210 Leu Asn Ala Leu Cys Lys Leu Ala Ala Val Leu Gly Ile Ser Ile Phe
211      690                    695                    700
213 aca tcc ttc gtg gga atc acc aag gct gca ccc atc ctc ttt gcc tca      2160
214 Thr Ser Phe Val Gly Ile Thr Lys Ala Ala Pro Ile Leu Phe Ala Ser
215 705                    710                    715                    720
217 gct gcc ctt gcc ctt ggc agc tct ctg gcc ctg aag ctg cct gag acc      2208
218 Ala Ala Leu Ala Leu Gly Ser Ser Leu Ala Leu Lys Leu Pro Glu Thr
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221 cgg ggg cag gtg ctg cag tga                                           2229
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223                740
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227 <211> LENGTH: 742
228 <212> TYPE: PRT
229 <213> ORGANISM: Homo sapiens
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238            20                    25                   30
241 Gly Leu Asp Arg Val Gln Asp Glu Tyr Ser Arg Arg Ser Tyr Ser Arg
242       35                    40                   45
245 Phe Glu Glu Glu Asp Asp Asp Asp Phe Pro Ala Pro Ser Asp Gly
246     50                    55                   60
249 Tyr Tyr Arg Gly Glu Gly Thr Gln Asp Glu Glu Glu Gly Gly Ala Ser
250 65                    70                   75                   80
253 Ser Asp Ala Thr Glu Gly His Asp Glu Asp Asp Glu Ile Tyr Glu Gly
254            85                    90                   95
257 Glu Tyr Gln Gly Ile Pro Arg Ala Glu Ser Gly Gly Lys Gly Glu Arg
258            100                   105                  110
261 Met Ala Asp Gly Ala Pro Leu Ala Gly Val Arg Gly Gly Leu Ser Asp
262       115                   120                  125
265 Gly Glu Gly Pro Pro Gly Gly Arg Gly Glu Ala Gln Arg Arg Lys Glu
266     130                   135                  140
269 Arg Glu Glu Leu Ala Gln Gln Tyr Glu Ala Ile Leu Arg Glu Cys Gly
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270 145                150                155                160
273 His Gly Arg Phe Gln Trp Thr Leu Tyr Phe Val Leu Gly Leu Ala Leu
274             165                170                175
277 Met Ala Asp Gly Val Glu Val Phe Val Val Gly Phe Val Leu Pro Ser
278             180                185                190
281 Ala Glu Lys Asp Met Cys Leu Ser Asp Ser Asn Lys Gly Met Leu Gly
282         195                200                205
285 Leu Ile Val Tyr Leu Gly Met Met Val Gly Ala Phe Leu Trp Gly Gly
286         210                215                220
289 Leu Ala Asp Arg Leu Gly Arg Arg Gln Cys Leu Leu Ile Ser Leu Ser
290 225                230                235                240
293 Val Asn Ser Val Phe Ala Phe Phe Ser Ser Phe Val Gln Gly Tyr Gly
294             245                250                255
297 Thr Phe Leu Phe Cys Arg Leu Leu Ser Gly Val Gly Ile Gly Gly Ser
298             260                265                270
301 Ile Pro Ile Val Phe Ser Tyr Phe Ser Glu Phe Leu Ala Gln Glu Lys
302         275                280                285
305 Arg Gly Glu His Leu Ser Trp Leu Cys Met Phe Trp Met Ile Gly Gly
306         290                295                300
309 Val Tyr Ala Ala Ala Met Ala Trp Ala Ile Ile Pro His Tyr Gly Trp
310 305                310                315                320
313 Ser Phe Gln Met Gly Ser Ala Tyr Gln Phe His Ser Trp Arg Val Phe
314             325                330                335
317 Val Leu Val Cys Ala Phe Pro Ser Val Phe Ala Ile Gly Ala Leu Thr
318             340                345                350
321 Thr Gln Pro Glu Ser Pro Arg Phe Phe Leu Glu Asn Gly Lys His Asp
322         355                360                365
325 Glu Ala Trp Met Val Leu Lys Gln Val His Asp Thr Asn Met Arg Ala
326         370                375                380
329 Lys Gly His Pro Glu Arg Val Phe Ser Val Thr His Ile Lys Thr Ile
330 385                390                395                400
333 His Gln Glu Asp Glu Leu Ile Glu Ile Gln Ser Asp Thr Gly Thr Trp
334             405                410                415
337 Tyr Gln Arg Trp Gly Val Arg Ala Leu Ser Leu Gly Gly Gln Val Trp
338         420                425                430
341 Gly Asn Phe Leu Ser Cys Phe Gly Pro Glu Tyr Arg Arg Ile Thr Leu
342         435                440                445
345 Met Met Met Gly Val Trp Phe Thr Met Ser Phe Ser Tyr Tyr Gly Leu
346         450                455                460
349 Thr Val Trp Phe Pro Asp Met Ile Arg His Leu Gln Ala Val Asp Tyr
350 465                470                475                480
353 Ala Ser Arg Thr Lys Val Phe Pro Gly Glu Arg Val Glu His Val Thr
354             485                490                495
357 Phe Asn Phe Thr Leu Glu Asn Gln Ile His Arg Gly Gly Gln Tyr Phe
358         500                505                510
361 Asn Asp Lys Phe Ile Gly Leu Arg Leu Lys Ser Val Ser Phe Glu Asp
362         515                520                525
365 Ser Leu Phe Glu Glu Cys Tyr Phe Glu Asp Val Thr Ser Ser Asn Thr
366         530                535                540
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VERIFICATION SUMMARY   DATE: 06/14/2005
PATENT APPLICATION: **US/10/537,512**   TIME: 09:05:03

Input Set : **A:\53529-5007-02-US Sequence Listing.txt**
Output Set: **N:\CRF4\06142005\J537512.raw**

L:13 M:270 C: Current Application Number differs, Replaced Current Application No
L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date